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IT IS CLAIMED:

- 1. A set of electrophoretic tag (e-tag) probes for detecting the binding of or interaction between each or any of a plurality of ligands and one or more target antiligands, the set comprising j members, and each of said e-tag probes having the form:
 - (D. Mi) L Ti, where
 - (a) D is a detection group comprising a detectable label;
 - (b) Ti is a ligand capable of binding to or interacting with a target antiligand,
- (c) L is a linking group connected to Ti by a bond that is cleavable by a selected cleaving agent when the probe is bound to or interacting with the target antiligand, wherein cleavage by said agent produces an e-tag reporter of the form (D, M_i) - L', where L' is the residue of Lattached to (D, Mi) after such cleavage,
- (d) Mi is a mobility modifier having a charge/mass ratio that imparts a unique and known electrophoretic mobility to a corresponding e-tag reporter of the form (D, Mi) - L', within a selected range of electrophoretic mobilities with respect to other e-tag reporters of the same form in the probe set; and
 - (e) (D, Mi)- includes both D Mi and Mi D -;
 - said uncleaved or partially cleaved e-tag probes, but not the corresponding e-tag reporter, having one or more chemical groups capable of reacting with or binding to a selected capture agent that is effective to
 - (i) impart a mobility to the probes bound to capture agent that prevents the probes from electrophoretically migrating within said range of electrophoretic mobilities or
 - (ii) immobilize the probes on a solid support.
- 2. The probe set of claim 1, for detecting each or any of a plurality of known, selected target nucleotide sequences, the set comprising j members, wherein:
- (a) Ti is an oligonucleotide target-binding moiety having a sequence of nucleotides U1 connected by intersubunit linkages Bi, i+1, where i includes all integers from 1 to n, and n is sufficient to allow the target-binding moiety to hybridize specifically with a target nucleotide sequence:
 - (b) L is a nucleotide joined to U1 in Ti through a nuclease-cleavable bond; and
- (c) each of the target-binding moieties contains at least one modification selected from the following:
 - (i) at least one nuclease-resistant bond Bi, i+1, where i includes at least 1;
 - (ii) U₁ containing a capture ligand capable of binding specifically to a capture agent; and
 - (iii) a nuclease-resistant bond Bi, +1, where i includes at least 1, and at least one nucleotide Ui containing a capture ligand capable of binding specifically to a capture agent, where $i \ge 1$.
- 3. The probe set of claim 1, wherein L includes at least a portion of an amino acid

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sequence that is recognized and cleaved by a selected peptidase.

4. The probe set of claim 1, wherein L includes at least a portion of an oligosaccharide that is recognized and cleaved by a selected hydrolytic enzyme.

5. The probe set of claim l , wherein L and T_j are linked by an ester linkage that is cleaved by a selected esterase.

- 6. The probe set of claim 1, wherein L and T_j are linked by a disulfide bond, and the antiligand is attached to an oxidase enzyme, such that in the presence of a substrate for the enzyme, H₂O₂ generated by the oxidase is effective to cleave the disulfide linkage in a probe bound to the antiligand.
 - 7. The probe of claim 1, wherein L and T_j are linked by a bond cleavable by singlet oxygen, wherein the antiligand is attached to a sensitizer capable of generating singlet oxygen when photoactivated.
 - The probe set of claim 1, for use in detecting the binding of each or any of a
 plurality of ligands to a target antiligand molecule, wherein the plurality of ligands are
 represented by T_j.

How is this limiting to Claim 1?

- 9. The probe set of claim 1, for use in screening for a ligand capable of binding to a receptor, wherein the ligands are represented by T_i and are members of a combinatorial library of small organic molecules, and the antiligand is the receptor. NOTE: This claim was originally written as a dependent claim to Claim 7 above.
- 10. The probe set of claim 1, for use in screening substrates of a selected enzyme antiligand, wherein the substrate comprises a fixed moiety L and a variable moiety T_j , and interaction of a substrate probe with the enzyme is effective to cleave the substrate to release the T_j moiety from the substrate.

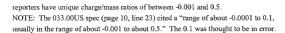
NOTE: This claim was originally written as a dependent claim to Claim 7 above.

- The probe set of claim 1, wherein each M_j has a unique charge/mass ratio by
 virtue of variations in mass, but not charge.
 - 12. The probe set of claim 1, wherein each M_j has a unique charge/mass ratio, by virtue of changes in both mass and charge.
- 40 13. The probe set of claim 12, containing at least 5 probes whose corresponding e-tag

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- 5 14. The probe set of claim 12, containing at least 9 probes whose corresponding e-tag reporters have unique charge/mass ratios of between -0.001 and 0.5.
 - 15. The probe set claim 12, wherein each M_j is formed of a selected number of negatively charged and/or positively charged amino acids.
 - 16. The probe set of claim 12, wherein each M_j includes an alkyl chain, and differs from other M_i in the set by 1-3 methylene groups in the chain.
 - 17. The probe set claim 1, wherein the detectable label is selected from the group consisting of a fluorophore, a chromophore, and an electrochemical compound capable of a detectable reaction in the presence of a redox agent.
 - 18. The probe set of claim 1, wherein the detectable label has a selected mass and charge.
 - 19. The probe set of claim 18, containing subsets of probes, each subset having a label with a unique mass/charge ratio.
 - 20. The probe set of claims 18 and 19, wherein the detectable label is a fluorophore.